

**AMENDMENTS TO THE CLAIMS**

1. (cancelled)

2. (previously presented) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein the 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residue from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted.

3. (previously presented) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein all of the 8<sup>th</sup> amino acid to 69<sup>th</sup> amino acid residues as measured from N terminal end are deleted, 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residues from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted.

4. (previously presented) An isolated polypeptide including the amino acid sequence described in SEQ ID NO:1 or 2.

5. (previously presented) An isolated DNA coding for the polypeptide of claim 2.

6. (cancelled)

7. (cancelled)

8. (previously presented) An isolated DNA coding for the polypeptide of claim 3.

9. (previously presented) An isolated DNA coding for the polypeptide of claim 4.

10. (previously presented) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein the 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residue from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted, wherein said polypeptide has membrane binding activity and induces Fas-mediated apoptotic activity.

11. (previously presented) An isolated polypeptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein all of the 8<sup>th</sup> amino acid to 69<sup>th</sup> amino acid residues as

measured from N terminal end are deleted, 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residues from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted, wherein said polypeptide has membrane binding activity and induces Fas-mediated apoptotic activity.

12. (currently amended) An isolated peptide having an amino acid sequence of natural human Fas ligand (SEQ ID NO:17) wherein at least four amino acid residues, including 128<sup>th</sup> and 131<sup>st</sup> amino acid residues are continuously deleted from the 111<sup>th</sup> amino acid to the 133<sup>rd</sup> amino acid residues as measured from N terminal end. ~~wherein the 129<sup>th</sup> amino acid and 130<sup>th</sup> amino acid residues as measured from N terminal end are both deleted, and at least one amino acid residue from 111<sup>th</sup> amino acid to 128<sup>th</sup> amino acid residues or at least one amino acid residue from 131<sup>st</sup> amino acid to 133<sup>rd</sup> amino acid residues as measured from N terminal end is deleted.~~